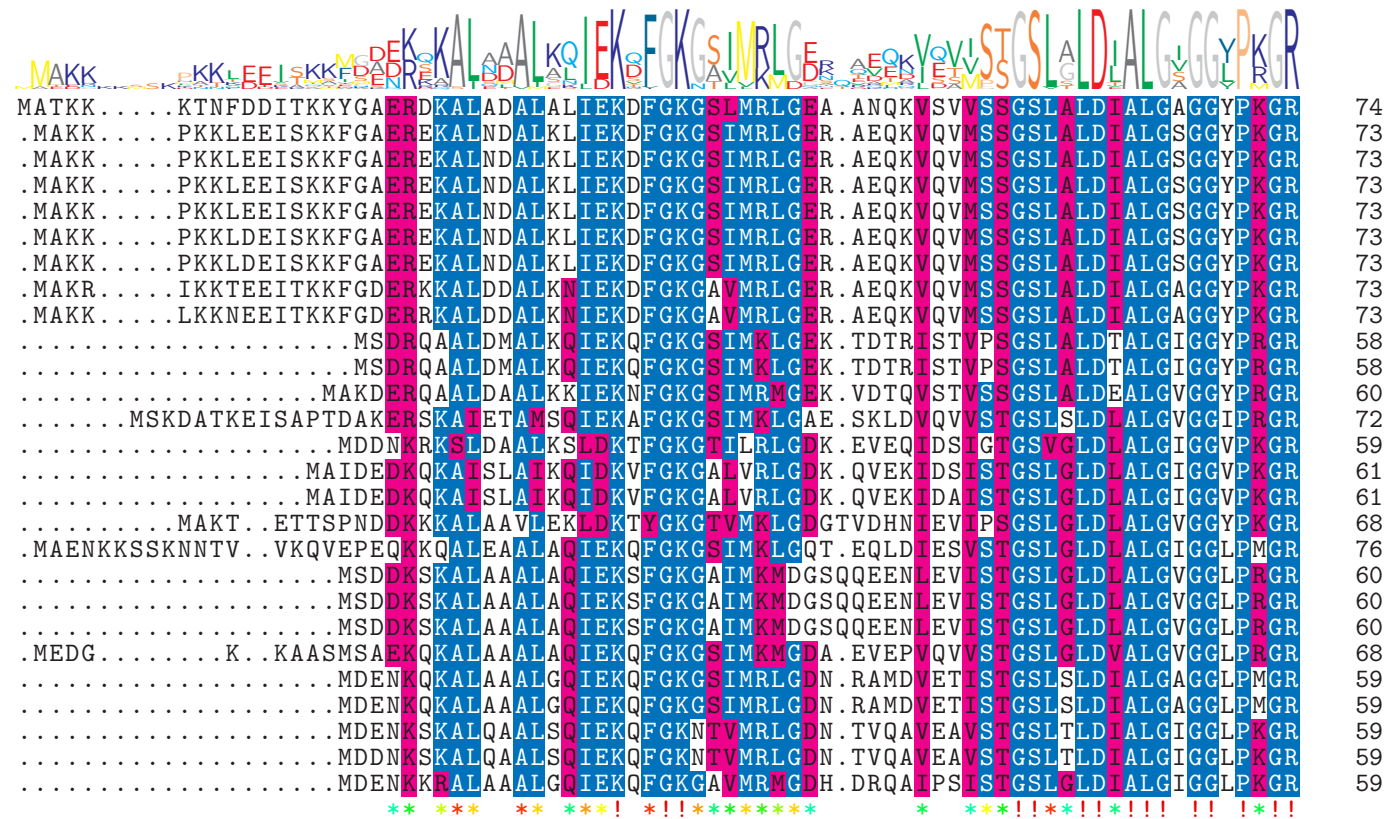


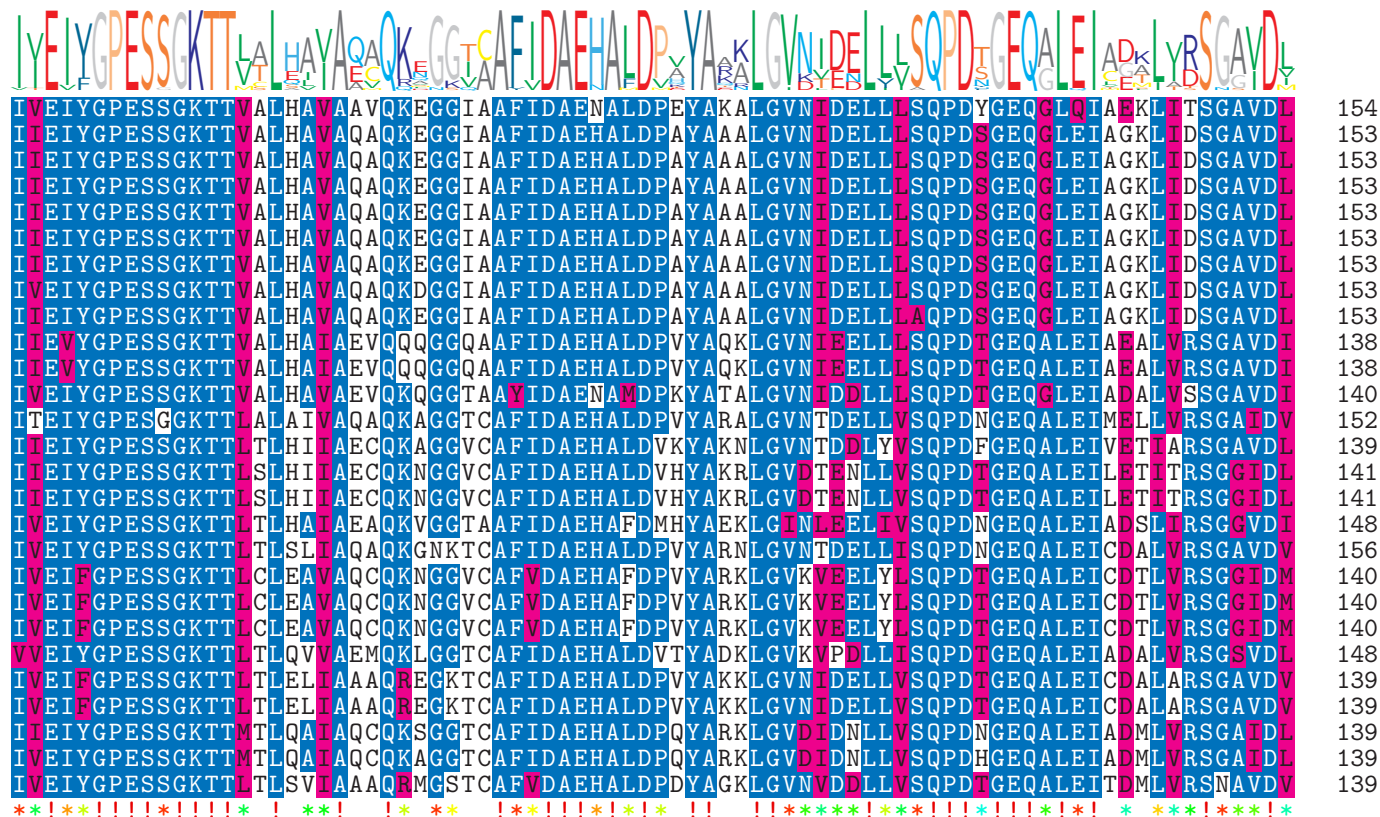
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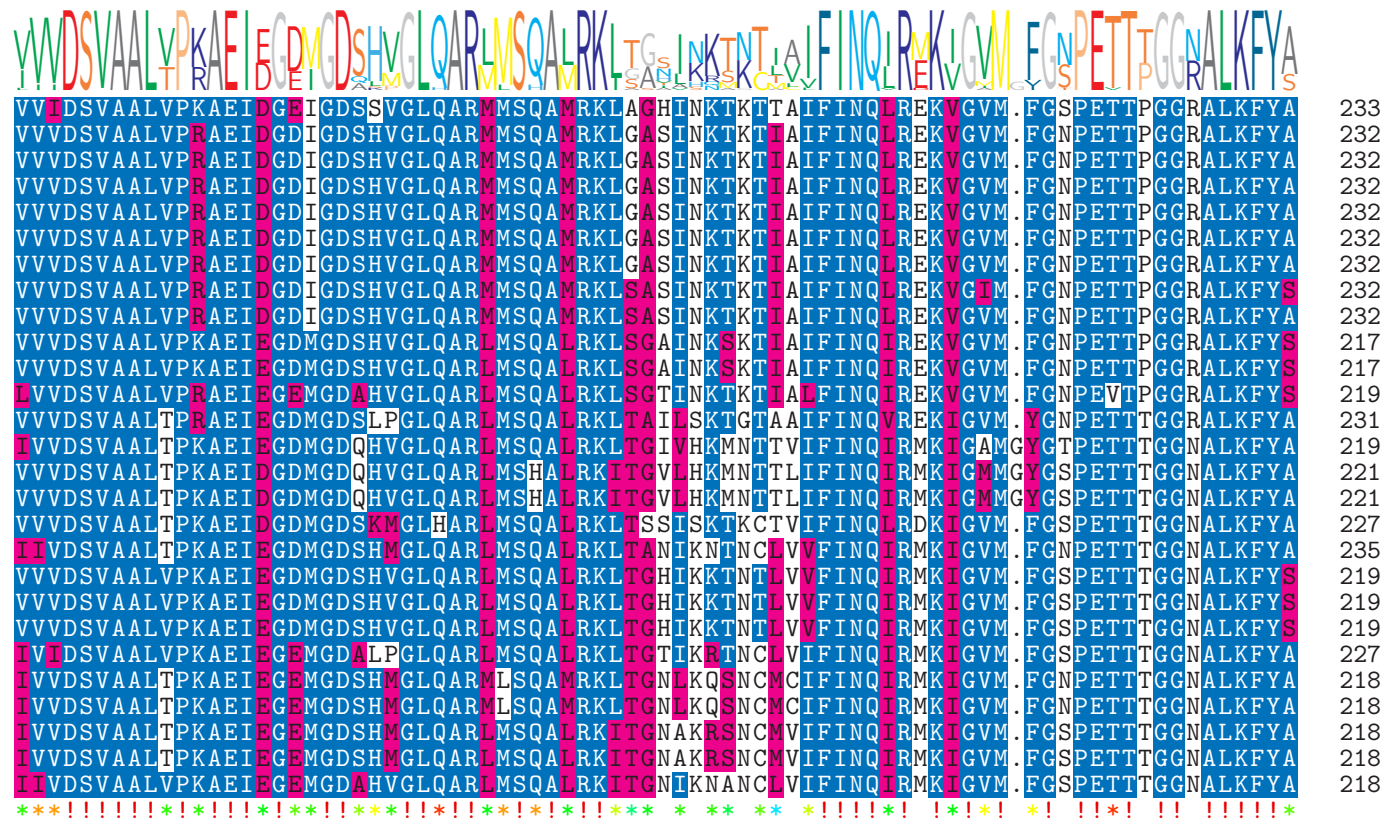
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consensus



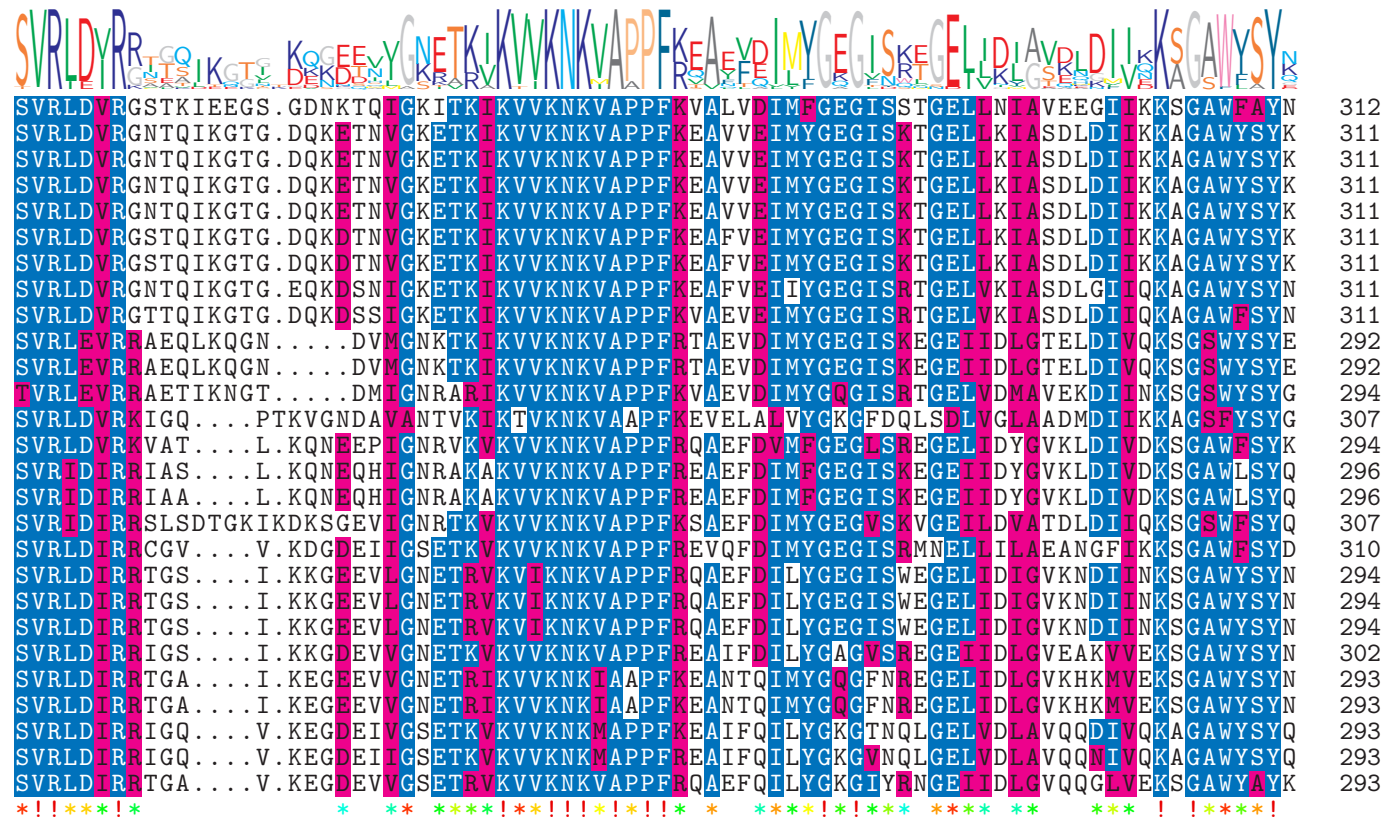
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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Sequence	Score
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DEKIGGGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IFDE ⁺ IDK ⁺ Q ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IFDE ⁺ IDK ⁺ Q ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IFDE ⁺ IDK ⁺ Q ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGGGS ⁺ ENAK ⁺ KYLAD ⁺ NPE ⁺ VFAE ⁺ IDH ⁺ Q ⁺ VRVQF ⁺ CL ⁺ IDGEEAAE....SKKDE....VAQVDS..VNEEVPLDLGDELEIE	381
DEKIGGGS ⁺ ENAK ⁺ KYLAD ⁺ NPE ⁺ IFDE ⁺ IDH ⁺ Q ⁺ VRVQF ⁺ CL ⁺ IDGEEIAE....SKKDE....AAQVDS..VNEEVTLDLGDELEIE	381
GEKIGGGS ⁺ ENAK ⁺ KFLAD ⁺ NPE ⁺ IFDD ⁺ IDHK ⁺ VRVQY ⁺ CL ⁺ IEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIEIE	382
GEKIGGGS ⁺ ENAK ⁺ KRYLAD ⁺ HQP ⁺ LFDE ⁺ IDRK ⁺ VRVKF ⁺ CL ⁺ LEESEEEESAMAVASE.....E.....TDDLALDLDNNGIEIE	377
EER ⁺ LG ⁺ GGREN ⁺ AKQ ⁺ FLKEN ⁺ KD ⁺ IMLM ⁺ IQEQ ⁺ IREHY ⁺ CL ⁺ DNNGVVQQQAETQEEL.EFE.E.....	348
EER ⁺ LG ⁺ GGREN ⁺ AKQ ⁺ FLKEN ⁺ KD ⁺ IMLM ⁺ IQEQ ⁺ IREHY ⁺ CL ⁺ DNNGVVQQQAETQEEL.EFE.E.....	348
SERIGGGREN ⁺ AKNYLAD ⁺ NHED ⁺ VEV ⁺ LRKV ⁺ RAAY ⁺ CL ⁺ ISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGK ⁺ EKTIAY ⁺ IAER ⁺ PEMEQ ⁺ EIRDRV ⁺ MAAIRAGNAGEAPALAPAPAAPEAAE.....	363
DKK ⁺ LG ⁺ GGRENS ⁺ KA ⁺ FLKEN ⁺ PEIA ⁺ DEIT ⁺ KA ⁺ IQNSM ⁺ CT ⁺ EGMI....SGSEDDE..GEE.....	343
DKK ⁺ LG ⁺ GGRENS ⁺ KA ⁺ LLKED ⁺ KALADEIT ⁺ TK ⁺ IKES ⁺ GSNEEI....MPLPDEP..LEEME.....	347
DKK ⁺ LG ⁺ GGRENS ⁺ KA ⁺ LLKED ⁺ KALADEIT ⁺ TK ⁺ IKES ⁺ GSNEEI....MPLPDEP..LEEME.....	347
DTKL ⁺ GQGRD ⁺ TVKEL ⁺ IKDN ⁺ PEIA ⁺ EELAK ⁺ IREKLQAK.....	343
GTKIGGK ⁺ NNAVK ⁺ WLKEN ⁺ PEIAD ⁺ KL ⁺ EQE ⁺ IRNTLA ⁺ SNPNS....VLKATASDSNDESD..VEEFSEEEEL.....	372
GAKIGGQ ⁺ KDNVR ⁺ VWLKEN ⁺ PEIS ⁺ DEID ⁺ AK ⁺ IRALNG ⁺ VEMHI....TEGTQDETD.....GERPEE.....	348
GAKIGGQ ⁺ KDNVR ⁺ VWLKEN ⁺ PEIS ⁺ DEID ⁺ AK ⁺ IRALNG ⁺ VEMHI....TEGTQDETD.....GERPEE.....	348
GAKIGGQ ⁺ KDNVR ⁺ VWLKEN ⁺ PEIS ⁺ DEID ⁺ AK ⁺ IRALNG ⁺ VEMHI....TEGTQDETD.....GERPEE.....	348
GERIGG ⁺ RDNCREF ⁺ LRN ⁺ EAL ⁺ AREI ⁺ ENK ⁺ VR ⁺ EHL ⁺ CVTPMG....AV.TLAEVEEED.....	352
GDKIGGK ⁺ KANACK ⁺ YLKEN ⁺ PEIA ⁺ KTLD ⁺ KKL ⁺ REMLLN ⁺ PENM....QLIAETSSAADDVE..FGAVPEEF.....	354
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GSKIGGK ⁺ KANA ⁺ AKYLE ⁺ ENPEI ⁺ GQIE ⁺ EQQ ⁺ IRAKLL ⁺ YTSAN....TKATPVAEDLADAD....I.....	347

logo

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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved